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Glu Val Pro Val Asn Phe Ala Glu Phe Ser Lys Lys Cys Ser Glu Arg 35 40 45

Trp Lys Thr Val Ser Gly Lys Glu Lys Ser Lys Phe Asp Glu Met Ala 50 55 60

Lys Ala Asp Lys Val Arg Tyr Asp Arg Glu Met Lys Asp Tyr Gly Pro 65 70 75 80

Ala Lys Gly Gly Lys Lys Lys Asp Pro Asn Ala Pro Lys Arg Pro
85 90 95

Pro Ser Gly Phe Phe Leu Phe Cys Ser Glu Phe Arg Pro Lys Ile Lys 100 105 110

Ser Thr Asn Pro Gly Ile Ser Ile Gly Asp Val Ala Lys Lys Leu Gly 115 120 125

Glu Met Trp Asn Asn Leu Asn Asp Ser Glu Lys Gln Pro Tyr Ile Thr 130 135 140

Lys Ala Ala Lys Leu Lys Glu Lys Tyr Glu Lys Asp Val Ala Asp Tyr 145 150 155 160

Lys Ser Lys Gly Lys Phe Asp Gly Ala Lys Gly Pro Ala Lys Val Ala 165 170 175

Arg Lys Lys Val Glu Glu Glu Asp Glu Glu Glu Glu Glu Glu Glu Glu Glu 180 185 190

Glu Glu Glu Glu Glu Asp Glu 195 200 <210> 325

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Ile Arg Phe Ser Asn Ile Ser Ala Ala Lys Ala Val Ala Asp Ala Ile 35 40 45

Arg Thr Ser Leu Gly Pro Lys Gly Met Asp Lys Met Ile Gln Asp Gly 50 55 60

Lys Gly Asp Val Thr Ile Thr Asn Asp Gly Ala Thr Ile Leu Lys Gln 65 70 75 80

Met Gln Val Leu His Pro Ala Ala Arg Met Leu Val Glu Leu Ser Lys 85 90 95

Ala Gln Asp Ile Glu Ala Gly Asp Gly Thr Thr Ser Val Val Ile Ile
100 105 110

Ala Gly Ser Leu Leu Asp Ser Cys Thr Lys Leu Leu Gln Lys Gly Ile 115 120 125

His Pro Thr Ile Ile Ser Glu Ser Phe Gln Lys Ala Leu Glu Lys Gly 130 135 140

Glu Thr Leu Leu Asn Ser Ala Thr Thr Ser Leu Asn Ser Lys Val Val
165 170 175

Ser Gln Tyr Ser Ser Leu Leu Ser Pro Met Ser Val Asn Ala Val Met 180 185 190

Lys Val Ile Asp Pro Ala Thr Ala Thr Ser Val Asp Leu Arg Asp Ile 195 200 205

Lys Ile Val Lys Lys Leu Gly Gly Thr Ile Asp Asp Cys Glu Leu Val 210 215 220

Glu Gly Leu Val Leu Thr Gln Lys Val Ser Asn Ser Gly Ile Thr Arg 225 230 235 240 Val Glu Lys Ala Lys Ile Gly Leu Ile Gln Phe Cys Leu Ser Ala Pro Lys Thr Asp Met Asp Asn Gln Ile Val Val Ser Asp Tyr Ala Gln Met 265 260 Asp Arg Val Leu Arg Glu Glu Arg Ala Tyr Ile Leu Asn Leu Val Lys 280 Gln Ile Lys Lys Thr Gly Cys Asn Val Leu Leu Ile Gln Lys Ser Ile 295 Leu Arg Asp Ala Leu Ser Asp Leu Ala Leu His Phe Leu Asn Lys Met 315 305 Lys Ile Met Val Ile Lys Asp Ile Glu Arg Glu Asp Ile Glu Phe Ile 330 Cys Lys Thr Ile Gly Thr Lys Pro Val Ala His Ile Asp Gln Phe Thr 345 Ala Asp Met Leu Gly Ser Ala Glu Leu Ala Glu Glu Val Asn Leu Asn Gly Ser Gly Lys Leu Leu Lys Ile Thr Gly Cys Ala Ser Pro Gly Lys 375 Thr Val Thr Ile Val Val Arg Gly Ser Asn Lys Leu Val Ile Glu Glu 385 390 Ala Glu Arg Ser Ile His Asp Ala Leu Cys Val Ile Arg Cys Leu Val 410 405 Lys Lys Arg Ala Leu Ile Ala Gly Gly Gly Ala Pro Glu Ile Glu Leu Ala Leu Arg Leu Thr Glu Tyr Ser Arg Thr Leu Ser Gly Met Glu Ser Tyr Cys Val Arg Ala Phe Ala Asp Ala Met Glu Val Ile Pro Ser Thr 455 Leu Ala Glu Asn Ala Gly Leu Asn Pro Ile Ser Thr Val Thr Glu Leu 475 470 465 Arg Asn Arg His Ala Gln Gly Glu Lys Thr Ala Gly Ile Asn Val Arg 490 Lys Gly Gly Ile Ser Asn Ile Leu Glu Glu Leu Val Val Gln Pro Leu 505 500 Leu Val Ser Val Ser Ala Leu Thr Leu Ala Thr Glu Thr Val Arg Ser 525 515 520

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<211> 144

<212> PRT

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Thr Ala Ala Leu Ile Phe Phe Ala Ile Trp His Ile Ile Ala Phe Asp 20 25 30

Glu Leu Lys Thr Asp Tyr Lys Asn Pro Ile Asp Gln Cys Asn Thr Leu 35 40 45

Asn Pro Leu Val Leu Pro Glu Tyr Leu Ile His Ala Phe Phe Cys Val 50 55 60

Met Phe Leu Cys Ala Ala Glu Trp Leu Thr Leu Gly Leu Asn Met Pro 65 70 75 80

Leu Leu Ala Tyr His Ile Trp Arg Tyr Met Ser Arg Pro Val Met Ser 85 90 95

Gly Pro Gly Leu Tyr Asp Pro Thr Thr Ile Met Asn Ala Asp Ile Leu 100 105 110

Ala Tyr Cys Gln Lys Glu Gly Trp Cys Lys Leu Ala Phe Tyr Leu Leu 115 120 125

Ala Phe Phe Tyr Tyr Leu Tyr Gly Met Ile Tyr Val Leu Val Ser Ser 130 135 140

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Glu Glu Leu Leu Lys Val Leu Gly Val Asn Val Met Leu Arg Lys Ile 20 25 30

Ala Val Ala Ala Ala Ser Lys Pro Ala Val Glu Ile Lys Gln Glu Gly 35 40 45

Asp Thr Phe Tyr Ile Lys Thr Ser Thr Thr Val Arg Thr Thr Glu Ile 50 55 60

Asn Phe Lys Val Gly Glu Glu Phe Glu Glu Gln Thr Val Asp Gly Arg
65 70 75 80

Pro Cys Lys Ser Leu Val Lys Trp Glu Ser Glu Asn Lys Met Val Cys 85 90 95

Glu Gln Lys Leu Leu Lys Gly Glu Gly Pro Lys Thr Ser Trp Thr Arg 100 105 110

Glu Leu Thr Asn Asp Gly Glu Leu Ile Leu Thr Met Thr Ala Asp Asp 115 120 125

Val Val Cys Thr Arg Val Tyr Val Arg Glu 130 135

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<211> 346

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Leu Gly Val Arg Gly Ala Pro Cys Glu Ala Val Arg Ile Pro Met Cys 20 25 30

Arg His Met Pro Trp Asn Ile Thr Arg Met Pro Asn His Leu His His 35 40 45

Ser Thr Gln Glu Asn Ala Ile Leu Ala Ile Glu Gln Tyr Glu Glu Leu 50 55 60

Val Asp Val Asn Cys Ser Ala Val Leu Arg Phe Phe Phe Cys Ala Met
65 70 75 80

Tyr Ala Pro Ile Cys Thr Leu Glu Phe Leu His Asp Pro Ile Lys Pro 85 90 95

Cys Lys Ser Val Cys Gln Arg Ala Arg Asp Asp Cys Glu Pro Leu Met 100 105 110

Lys Met Tyr Asn His Ser Trp Pro Glu Ser Leu Ala Cys Asp Glu Leu 115 120 125

Pro Val Tyr Asp Arg Gly Val Cys Ile Ser Pro Glu Ala Ile Val Thr 130 135 140

Asp Leu Pro Glu Asp Val Lys Trp Ile Asp Ile Thr Pro Asp Met Met 145 150 155 160

Val Gln Glu Arg Pro Leu Asp Val Asp Cys Lys Arg Leu Ser Pro Asp 165 Arg Cys Lys Cys Lys Lys Val Lys Pro Thr Leu Ala Thr Tyr Leu Ser 180 Lys Asn Tyr Ser Tyr Val Ile His Ala Lys Ile Lys Ala Val Gln Arg 200 Ser Gly Cys Asn Glu Val Thr Thr Val Val Asp Val Lys Glu Ile Phe 215 Lys Ser Ser Ser Pro Ile Pro Arg Thr Gln Val Pro Leu Ile Thr Asn 235 230 Ser Ser Cys Gln Cys Pro His Ile Leu Pro His Gln Asp Val Leu Ile 250 245 Met Cys Tyr Glu Trp Arg Ser Arg Met Met Leu Leu Glu Asn Cys Leu 260 Val Glu Lys Trp Arg Asp Gln Leu Ser Lys Arg Ser Ile Gln Trp Glu 280 Glu Arg Leu Gln Glu Gln Arg Arg Thr Val Gln Asp Lys Lys Thr 295 Ala Gly Arg Thr Ser Arg Ser Asn Pro Pro Lys Pro Lys Gly Lys Pro 310 305 Pro Ala Pro Lys Pro Ala Ser Pro Lys Lys Asn Ile Lys Thr Arg Ser 330 325 Ala Gln Lys Arg Thr Asn Pro Lys Arg Val

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Arg Arg Lys Glu Lys Ser Arg Asp Ala Ala Arg Ser Arg Arg Ser Lys 20 25 30

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35 40 45

Asn Val Ser Ser His Leu Asp Lys Ala Ser Val Met Arg Leu Thr Ile

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Asp	Gly	Phe	Val 100	Met	Val	Leu	Thr	Asp 105	Asp	Gly	Asp	Met	Ile 110	Tyr	Ile
Ser	Asp	Asn 115	Val	Asn	Lys	Tyr	Met 120	Gly	Leu	Thr	Gln	Phe 125	Glu	Leu	Thr
Gly	His 130	Ser	Val	Phe	Asp	Phe 135	Thr	His	Pro	Cys	Asp 140	His	Glu	Glu	Met
Arg 145	Glu	Met	Leu	Thr	His 150	Arg	Asn	Gly	Leu	Val 155	Lys	Lys	Gly	Lys	Glu 160
Gln	Asn	Thr	Gln	Arg 165	Ser	Phe	Phe	Leu	Arg 170	Met	Lys	Cys	Thr	Leu 175	Thr
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His	Cys	Thr 195	Gly	His	Ile	His	Val 200	Tyr	Asp	Thr	Asn	Ser 205	Asn	Gln	Pro
Gln	Cys 210	Gly	Tyr	Lys	Lys	Pro 215	Pro	Met	Thr	Cys	Leu 220	Val	Leu	Ile	Cys
Glu 225	Pro	Ile	Pro	His	Pro 230	Ser	Asn	Ile	Glu	Ile 235	Pro	Leu	Asp	Ser	Lys 240
Thr	Phe	Leu	Ser	Arg 245	His	Ser	Leu	Asp	Met 250	Lys	Phe	Ser	Tyr	Cys 255	Asp
Glu	Arg	Ile	Thr 260		Leu	Met	Gly	Tyr 265	Glu	Pro	Glu	Glu	Leu 270	Leu	Gly
Arg	Ser	Ile 275	Tyr	Glu	Tyr	Tyr	His 280	Ala	Leu	Asp	Ser	Asp 285	His	Leu	Thr
Lys	Thr 290	His	His	Asp	Met	Phe 295	Thr	Lys	Gly	Gln	Val 300	Thr	Thr	Gly	Gln
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Ala	Thr	Val	Ile	Tyr 325	Asn	Thr	Lys	Asn	Ser 330	Gln	Pro	Gln	Cys	Ile 335	Val
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345 350 340 Ser Leu Gln Gln Thr Glu Cys Val Leu Lys Pro Val Glu Ser Ser Asp 360 Met Lys Met Thr Gln Leu Phe Thr Lys Val Glu Ser Glu Asp Thr Ser 375 Ser Leu Phe Asp Lys Leu Lys Lys Glu Pro Asp Ala Leu Thr Leu Leu 390 395 Ala Pro Ala Ala Gly Asp Thr Ile Ile Ser Leu Asp Phe Gly Ser Asn 405 Asp Thr Glu Thr Asp Asp Gln Gln Leu Glu Glu Val Pro Leu Tyr Asn 425 Asp Val Met Leu Pro Ser Pro Asn Glu Lys Leu Gln Asn Ile Asn Leu Ala Met Ser Pro Leu Pro Thr Ala Glu Thr Pro Lys Pro Leu Arg Ser 455 Ser Ala Asp Pro Ala Leu Asn Gln Glu Val Ala Leu Lys Leu Glu Pro 475 Asn Pro Glu Ser Leu Glu Leu Ser Phe Thr Met Pro Gln Ile Gln Asp 490 Gln Thr Pro Ser Pro Ser Asp Gly Ser Thr Arg Gln Ser Ser Pro Glu 505 Pro Asn Ser Pro Ser Glu Tyr Cys Phe Tyr Val Asp Ser Asp Met Val 515 Asn Glu Phe Lys Leu Glu Leu Val Glu Lys Leu Phe Ala Glu Asp Thr 535 Glu Ala Lys Asn Pro Phe Ser Thr Gln Asp Thr Asp Leu Asp Leu Glu 555 Met Leu Ala Pro Tyr Ile Pro Met Asp Asp Phe Gln Leu Arg Ser Phe Asp Gln Leu Ser Pro Leu Glu Ser Ser Ser Ala Ser Pro Glu Ser 585 Ala Ser Pro Gln Ser Thr Val Thr Val Phe Gln Gln Thr Gln Ile Gln 595 Glu Pro Thr Ala Asn Ala Thr Thr Thr Thr Ala Thr Thr Asp Glu Leu Lys Thr Val Thr Lys Asp Arg Met Glu Asp Ile Lys Ile Leu Ile Ala

635 640 630 625 Ser Pro Ser Pro Thr His Ile His Lys Glu Thr Thr Ser Ala Thr Ser 645 Ser Pro Tyr Arg Asp Thr Gln Ser Arg Thr Ala Ser Pro Asn Arg Ala 665 Gly Lys Gly Val Ile Glu Gln Thr Glu Lys Ser His Pro Arg Ser Pro 675 Asn Val Leu Ser Val Ala Leu Ser Gln Arg Thr Thr Val Pro Glu Glu Glu Leu Asn Pro Lys Ile Leu Ala Leu Gln Asn Ala Gln Arg Lys Arg 715 Lys Met Glu His Asp Gly Ser Leu Phe Gln Ala Val Gly Ile Gly Thr 730 725 Leu Leu Gln Gln Pro Asp Asp His Ala Ala Thr Thr Ser Leu Ser Trp 745 Lys Arg Val Lys Gly Cys Lys Ser Ser Glu Gln Asn Gly Met Glu Gln Lys Thr Ile Ile Leu Ile Pro Ser Asp Leu Ala Cys Arg Leu Leu Gly 775 Gln Ser Met Asp Glu Ser Gly Leu Pro Gln Leu Thr Ser Tyr Asp Cys 795 Glu Val Asn Ala Pro Ile Gln Gly Ser Arg Asn Leu Leu Gln Gly Glu 810 Glu Leu Leu Arg Ala Leu Asp Gln Val Asn 820 <210> 331 <211> 92 <212> PRT <213> Homo sapiens <400> 331 Met Ala Tyr Arg Gly Gln Gly Gln Lys Val Gln Lys Val Met Val Gln Pro Ile Asn Leu Ile Phe Arg Tyr Leu Gln Asn Arg Ser Arg Ile Gln Val Trp Leu Tyr Glu Gln Val Asn Met Arg Ile Glu Gly Cys Ile Ile Gly Phe Asp Glu Tyr Met Asn Leu Val Leu Asp Asp Ala Glu Glu Ile 50 55 60

His Ser Lys Thr Lys Ser Arg Lys Gln Leu Gly Arg Ile Met Leu Lys 65 70 75 80

Gly Asp Asn Ile Thr Leu Leu Gln Ser Val Ser Asn \$85\$ 90

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<211> 235

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<400> 332

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5 10 15

Thr Glu Ala Ala Leu Gly Asp Ala Ala Gln Glu Pro Thr Gly Asn Asn 20 25 30

Ala Glu Ile Cys Leu Leu Pro Leu Asp Tyr Gly Pro Cys Arg Ala Leu 35 40 45

Leu Leu Arg Tyr Tyr Tyr Asp Arg Tyr Thr Gln Ser Cys Arg Gln Phe 50 55 60

Leu Tyr Gly Gly Cys Glu Gly Asn Ala Asn Asn Phe Tyr Thr Trp Glu 65 70 75 80

Ala Cys Asp Asp Ala Cys Trp Arg Ile Glu Lys Val Pro Lys Val Cys 85 90 95

Arg Leu Gln Val Ser Val Asp Asp Gln Cys Glu Gly Ser Thr Glu Lys
100 105 110

Tyr Phe Phe Asn Leu Ser Ser Met Thr Cys Glu Lys Phe Phe Ser Gly 115 120 125

Gly Cys His Arg Asn Arg Ile Glu Asn Arg Phe Pro Asp Glu Ala Thr

Cys Met Gly Phe Cys Ala Pro Lys Lys Ile Pro Ser Phe Cys Tyr Ser 145 150 155 160

Pro Lys Asp Glu Gly Leu Cys Ser Ala Asn Val Thr Arg Tyr Tyr Phe 165 170 175

Asn Pro Arg Tyr Arg Thr Cys Asp Ala Phe Thr Tyr Thr Gly Cys Gly
180 185 190

Gly Asn Asp Asn Asn Phe Val Ser Arg Glu Asp Cys Lys Arg Ala Cys 195 200 205 Ala Lys Ala Leu Lys Lys Lys Lys Met Pro Lys Leu Arg Phe Ala 210 215 220

Ser Arg Ile Arg Lys Ile Arg Lys Lys Gln Phe 225 230 235

<210> 333

<211> 291

<212> PRT

<213> Homo sapiens

<400> 333

Met Gln Arg Ala Arg Pro Thr Leu Trp Ala Ala Ala Leu Thr Leu Leu
5 10 15

Val Leu Leu Arg Gly Pro Pro Val Ala Arg Ala Gly Ala Ser Ser Gly 20 25 30

Gly Leu Gly Pro Val Val Arg Cys Glu Pro Cys Asp Ala Arg Ala Leu 35 40 45

Ala Gln Cys Ala Pro Pro Pro Ala Val Cys Ala Glu Leu Val Arg Glu
50 55 60

Pro Gly Cys Gly Cys Cys Leu Thr Cys Ala Leu Ser Glu Gly Gln Pro 65 70 75 80

Cys Gly Ile Tyr Thr Glu Arg Cys Gly Ser Gly Leu Arg Cys Gln Pro 85 90 95

Ser Pro Asp Glu Ala Arg Pro Leu Gln Ala Leu Leu Asp Gly Arg Gly
100 105 110

Leu Cys Val Asn Ala Ser Ala Val Ser Arg Leu Arg Ala Tyr Leu Leu 115 120 125

Pro Ala Pro Pro Ala Pro Gly Asn Ala Ser Glu Ser Glu Glu Asp Arg 130 135 140

Ser Ala Gly Ser Val Glu Ser Pro Ser Val Ser Ser Thr His Arg Val 145 150 155 160

Ser Asp Pro Lys Phe His Pro Leu His Ser Lys Ile Ile Ile Ile Lys 165 170 175

Lys Gly His Ala Lys Asp Ser Gln Arg Tyr Lys Val Asp Tyr Glu Ser 180 185 190

Gln Ser Thr Asp Thr Gln Asn Phe Ser Ser Glu Ser Lys Arg Glu Thr 195 200 205

Glu Tyr Gly Pro Cys Arg Glu Met Glu Asp Thr Leu Asn His Leu 210 215 220

Lys Phe Leu Asn Val Leu Ser Pro Arg Gly Val His Ile Pro Asn Cys 225 230 235 240

Asp Lys Lys Gly Phe Tyr Lys Lys Lys Gln Cys Arg Pro Ser Lys Gly
245 250 255

Arg Lys Arg Gly Phe Cys Trp Cys Val Asp Lys Tyr Gly Gln Pro Leu 260 265 270

Pro Gly Tyr Thr Thr Lys Gly Lys Glu Asp Val His Cys Tyr Ser Met 275 280 285

Gln Ser Lys 290

<210> 334

<211> 582

<212> PRT

<213> Homo sapiens

<400> 334

Glu Ser Lys Gly Ala Ser Ser Cys Arg Leu Leu Phe Cys Leu Leu Ile 5 10 15

Ser Ala Thr Val Phe Arg Pro Gly Leu Gly Trp Tyr Thr Val Asn Ser 20 25 30

Ala Tyr Gly Asp Thr Ile Ile Ile Pro Cys Arg Leu Asp Val Pro Gln
35 40 45

Asn Leu Met Phe Gly Lys Trp Lys Tyr Glu Lys Pro Asp Gly Ser Pro 50 55 60

Val Phe Ile Ala Phe Arg Ser Ser Thr Lys Lys Ser Val Gln Tyr Asp 65 70 75 80

Asp Val Pro Glu Tyr Lys Asp Arg Leu Asn Leu Ser Glu Asn Tyr Thr 85 90 95

Leu Ser Ile Ser Asn Ala Arg Ile Ser Asp Glu Lys Arg Phe Val Cys
100 105 110

Met Leu Val Thr Glu Asp Asn Val Phe Glu Ala Pro Thr Ile Val Lys 115 120 125

Val Phe Lys Gln Pro Ser Lys Pro Glu Ile Val Ser Lys Ala Leu Phe 130 135 140

Leu Glu Thr Glu Gln Leu Lys Lys Leu Gly Asp Cys Ile Ser Glu Asp 145 150 155 160

Ser Tyr Pro Asp Gly Asn Ile Thr Trp Tyr Arg Asn Gly Lys Val Leu

				165					170					175	
His	Pro	Leu	Glu 180	Gly	Ala	Val	Val	Ile 185	Ile	Phe	Lys	Lys	Glu 190	Met	Asp
Pro	Val	Thr 195	Gln	Leu	Tyr	Thr	Met 200	Thr	Ser	Thr	Leu	Glu 205	Tyr	Lys	Thr
Thr	Lys 210	Ala	Asp	Ile	Gln	Met 215	Pro	Phe	Thr	Cys	Ser 220	Val	Thr	Tyr	Tyr
Gly 225	Pro	Ser	Gly	Gln	Lys 230	Thr	Ile	His	Ser	Glu 235	Gln	Ala	Val	Phe	Asp 240
Ile	Tyr	Tyr	Pro	Thr 245	Glu	Gln	Val	Thr	Ile 250	Gln	Val	Leu	Pro	Pro 255	Lys
Asn	Ala	Ile	Lys 260	Glu	Gly	Asp	Asn	Ile 265	Thr	Leu	Lys	Cys	Leu 270	Gly	Asn
Gly	Asn	Pro 275	Pro	Pro	Glu	Glu	Phe 280	Leu	Phe	Tyr	Leu	Pro 285	Gly	Gln	Pro
Glu	Gly 290	Ile	Arg	Ser	Ser	Asn 295	Thr	Tyr	Thr	Leu	Thr 300	Asp	Val	Arg	Arg
Asn 305	Ala	Thr	Gly	Asp	Tyr 310	Lys	Cys	Ser	Leu	Ile 315	Asp	Lys	Lys	Ser	Met 320
Ile	Ala	Ser	Thr	Ala 325	Ile	Thr	Val	His	Tyr 330	Leu	Asp	Leu	Ser	Leu 335	Asn
Pro	Ser	Gly	Glu 340	Val	Thr	Arg	Gln	Ile 345	Gly	Asp	Ala	Leu	Pro 350	Val	Ser
Cys	Thr	Ile 355	Ser	Ala	Ser	Arg	Asn 360	Ala	Thr	Val	Val	Trp 365	Met	Lys	Asp
Asn	Ile 370		Leu	Arg	Ser	Ser 375	Pro	Ser	Phe	Ser	Ser 380	Leu	His	Tyr	Gln
Asp 385	Ala	Gly	Asn	Tyr	Val 390	Cys	Glu	Thr	Ala	Leu 395	Gln	Glu	Val	Glu	Gly 400
Leu	Lys	Lys	Arg	Glu 405	Ser	Leu	Thr	Leu	Ile 410	Val	Glu	Gly	Lys	Pro 415	Gln
Ile	Lys	Met	Thr 420	Lys	Lys	Thr	Asp	Pro 425	Ser	Gly	Leu	Ser	Lys 430	Thr	Ile
Ile	Cys	His 435	Val	Glu	Gly	Phe	Pro 440	Lys	Pro	Ala	Ile	Gln 445		Thr	Ile
Thr	Glv	Ser	Glv	Ser	Val	Ile	Asn	Gln	Thr	Glu	Glu	Ser	Pro	Tyr	Ile

460 455 450 Asn Gly Arg Tyr Tyr Ser Lys Ile Ile Ile Ser Pro Glu Glu Asn Val Thr Leu Thr Cys Thr Ala Glu Asn Gln Leu Glu Arg Thr Val Asn Ser 490 Leu Asn Val Ser Ala Ile Ser Ile Pro Glu His Asp Glu Ala Asp Glu 505 Ile Ser Asp Glu Asn Arg Glu Lys Val Asn Asp Gln Ala Lys Leu Ile 515 Val Gly Ile Val Val Gly Leu Leu Leu Ala Ala Leu Val Ala Gly Val 535 Val Tyr Trp Leu Tyr Met Lys Lys Ser Lys Thr Ala Ser Lys His Val 545 Asn Lys Asp Leu Gly Asn Met Glu Glu Asn Lys Lys Leu Glu Glu Asn 570 565 Asn His Lys Thr Glu Ala 580 <210> 335 <211> 709 <212> PRT <213> Homo sapiens <400> 335 Met Ala Glu Val Glu Asp Gln Ala Ala Arg Asp Met Lys Arg Leu Glu Glu Lys Asp Lys Glu Arg Lys Asn Val Lys Gly Ile Arg Asp Asp Ile 20 Glu Glu Glu Asp Asp Gln Glu Ala Tyr Phe Arg Tyr Met Ala Glu Asn Pro Thr Ala Gly Val Val Glu Glu Glu Glu Glu Asp Asn Leu Glu Tyr Asp Ser Asp Gly Asn Pro Ile Ala Pro Thr Lys Lys Ile Ile Asp Pro 65 Leu Pro Pro Ile Asp His Ser Glu Ile Asp Tyr Pro Pro Phe Glu Lys Asn Phe Tyr Asn Glu His Glu Glu Ile Thr Asn Leu Thr Pro Gln Gln 105

Leu Ile Asp Leu Arg His Lys Leu Asn Leu Arg Val Ser Gly Ala Ala 120 Pro Pro Arg Pro Gly Ser Ser Phe Ala His Phe Gly Phe Asp Glu Gln 135 Leu Met His Gln Ile Arg Lys Ser Glu Tyr Thr Gln Pro Thr Pro Ile 155 150 Gln Cys Gln Gly Val Pro Val Ala Leu Ser Gly Arg Asp Met Ile Gly 165 Ile Ala Lys Thr Gly Ser Gly Lys Thr Ala Ala Phe Ile Trp Pro Met 185 Leu Ile His Ile Met Asp Gln Lys Glu Leu Glu Pro Gly Asp Gly Pro Ile Ala Val Ile Val Cys Pro Thr Arg Glu Leu Cys Gln Gln Ile His 215 Ala Glu Cys Lys Arg Phe Gly Lys Ala Tyr Asn Leu Arg Ser Val Ala 235 Val Tyr Gly Gly Ser Met Trp Glu Gln Ala Lys Ala Leu Gln Glu 245 Gly Ala Glu Ile Val Val Cys Thr Pro Gly Arg Leu Ile Asp His Val 265 Lys Lys Lys Ala Thr Asn Leu Gln Arg Val Ser Tyr Leu Val Phe Asp 275 Glu Ala Asp Arg Met Phe Asp Met Gly Phe Glu Tyr Gln Val Arg Ser 295 Ile Ala Ser His Val Arg Pro Asp Arg Gln Thr Leu Leu Phe Ser Ala 315 Thr Phe Arg Lys Lys Ile Glu Lys Leu Ala Arg Asp Ile Leu Ile Asp Pro Ile Arg Val Val Gln Gly Asp Ile Gly Glu Ala Asn Glu Asp Val 345 Thr Gln Ile Val Glu Ile Leu His Ser Gly Pro Ser Lys Trp Asn Trp 355 Leu Thr Arg Arg Leu Val Glu Phe Thr Ser Ser Gly Ser Val Leu Leu 375 Phe Val Thr Lys Lys Ala Asn Ala Glu Glu Leu Ala Asn Asn Leu Lys 400 395

Gln Glu Gly His Asn Leu Gly Leu Leu His Gly Asp Met Asp Gln Ser 410 Glu Arg Asn Lys Val Ile Ser Asp Phe Lys Lys Asp Ile Pro Val 425 Leu Val Ala Thr Asp Val Ala Ala Arg Gly Leu Asp Ile Pro Ser Ile Lys Thr Val Ile Asn Tyr Asp Val Ala Arg Asp Ile Asp Thr His Thr 455 His Arg Ile Gly Arg Thr Gly Arg Ala Gly Glu Lys Gly Val Ala Tyr 470 475 Thr Leu Leu Thr Pro Lys Asp Ser Asn Phe Ala Gly Asp Leu Val Arg 485 490 Asn Leu Glu Gly Ala Asn Gln His Val Ser Lys Glu Leu Leu Asp Leu 505 Ala Met Gln Asn Ala Trp Phe Arg Lys Ser Arg Phe Lys Gly Gly Lys 520 515 Gly Lys Lys Leu Asn Ile Gly Gly Gly Leu Gly Tyr Arg Glu Arg Pro Gly Leu Gly Ser Glu Asn Met Asp Arg Gly Asn Asn Asn Val Met 555 Ser Asn Tyr Glu Ala Tyr Lys Pro Ser Thr Gly Ala Met Gly Asp Arg 565 Leu Thr Ala Met Lys Ala Ala Phe Gln Ser Gln Tyr Lys Ser His Phe 585 Val Ala Ser Leu Ser Asn Gln Lys Ala Gly Ser Ser Ala Ala Gly 595 Ala Ser Gly Trp Thr Ser Ala Gly Ser Leu Asn Ser Val Pro Thr Asn Ser Ala Gln Gln Gly His Asn Ser Pro Asp Ser Pro Val Thr Ser Ala 635 Ala Lys Gly Ile Pro Gly Phe Gly Asn Thr Gly Asn Ile Ser Gly Ala 645 Pro Val Thr Tyr Pro Ser Ala Gly Ala Gln Gly Val Asn Asn Thr Ala 665 Ser Gly Asn Asn Ser Arg Glu Gly Thr Gly Gly Ser Asn Gly Lys Arg 680

Glu Arg Tyr Thr Glu Asn Arg Gly Ser Ser Pro Ser Gln Ser Arg Arg 690 695 700

Asp Trp Gln Ser Ala 705

<210> 336

<211> 480

<212> PRT

<213> Homo sapiens

<400> 336

Met Ile Arg Ala Ala Pro Pro Pro Leu Phe Leu Leu Leu Leu Leu Leu 15

Leu Leu Val Ser Trp Ala Ser Arg Gly Glu Ala Ala Pro Asp Gln 20 25 30

Asp Glu Ile Gln Arg Leu Pro Gly Leu Ala Lys Gln Pro Ser Phe Arg
35 40 45

Gln Tyr Ser Gly Tyr Leu Lys Ser Ser Gly Ser Lys His Leu His Tyr 50 55 60

Trp Phe Val Glu Ser Gln Lys Asp Pro Glu Asn Ser Pro Val Val Leu 65 70 75 80

Trp Leu Asn Gly Gly Pro Gly Cys Ser Ser Leu Asp Gly Leu Leu Thr
85 90 95

Glu His Gly Pro Phe Leu Val Gln Pro Asp Gly Val Thr Leu Glu Tyr 100 105 110

Asn Pro Tyr Ser Trp Asn Leu Ile Ala Asn Val Leu Tyr Leu Glu Ser 115 120 125

Pro Ala Gly Val Gly Phe Ser Tyr Ser Asp Asp Lys Phe Tyr Ala Thr 130 135 140

Asn Asp Thr Glu Val Ala Gln Ser Asn Phe Glu Ala Leu Gln Asp Phe 145 150 155 160

Phe Arg Leu Phe Pro Glu Tyr Lys Asn Asn Lys Leu Phe Leu Thr Gly 165 170 175

Glu Ser Tyr Ala Gly Ile Tyr Ile Pro Thr Leu Ala Val Leu Val Met 180 185 190

Gln Asp Pro Ser Met Asn Leu Gln Gly Leu Ala Val Gly Asn Gly Leu 195 200 205

Ser Ser Tyr Glu Gln Asn Asp Asn Ser Leu Val Tyr Phe Ala Tyr Tyr 210 215 220

His 225	Gly	Leu	Leu	Gly	Asn 230	Arg	Leu	Trp	Ser	Ser 235	Leu	Gln	Thr	His	Cys 240
Cys	Ser	Gln	Asn	Lys 245	Cys	Asn	Phe	Tyr	Asp 250	Asn	Lys	Asp	Leu	Glu 255	Cys
Val	Thr	Asn	Leu 260	Gln	Glu	Val	Ala	Arg 265	Ile	Val	Gly	Asn	Ser 270	Gly	Leu
Asn	Ile	Tyr 275	Asn	Leu	Tyr	Ala	Pro 280	Cys	Ala	Gly	Gly	Val 285	Pro	Ser	His
Phe	Arg 290	Tyr	Glu	Lys	Āsp	Thr 295	Val	Vāl	Val	Gln	Asp 300	Leu	Gly	Asn	Ile
Phe 305	Thr	Arg	Leu	Pro	Leu 310	Lys	Arg	Met	Trp	His 315	Gln	Ala	Leu	Leu	Arg 320
Ser	Gly	Asp	Lys	Val 325	Arg	Met	Asp	Pro	Pro 330	Cys	Thr	Asn	Thr	Thr 335	Ala
Ala	Ser	Thr	Tyr 340	Leu	Asn	Asn	Pro	Tyr 345	Val	Arg	Lys	Ala	Leu 350	Asn	Ile
Pro	Glu	Gln 355	Leu	Pro	Gln	Trp	Asp 360	Met	Cys	Asn	Phe	Leu 365	Val	Asn	Leu
Gln	Tyr 370	Arg	Arg	Leu	Tyr	Arg 375	Ser	Met	Asn	Ser	Gln 380	Tyr	Leu	Lys	Leu
Leu 385	Ser	Ser	Gln	Lys	Tyr 390	Gln	Ile	Leu	Leu	Tyr 395	Asn	Gly	Asp	Val	Asp 400
Met	Ala	Cys	Asn	Phe 405	Met	Gly	Asp	Glu	Trp 410	Phe	Val	Asp	Ser	Leu 415	Asn
Gln	Lys	Met	Glu 420	Val	Gln	Arg	Arg	Pro 425	Trp	Leu	Val	Lys	Tyr 430	Gly	Asp
Ser	Gly	Glu 435	Gln	Ile	Ala	Gly	Phe 440	Val	Lys	Glu	Phe	Ser 445	His	Ile	Ala
Phe	Leu 450	Thr	Ile	Lys	Gly	Ala 455	Gly	His	Met	Val	Pro 460	Thr	Asp	Lys	Pro
Leu 465	Ala	Ala	Phe	Thr	Met 470	Phe	Ser	Arg	Phe	Leu 475	Asn	Lys	Gln	Pro	Tyr 480

<210> 337

<211> 543 <212> PRT <213> Homo sapiens

<400> 337 Met Ala Ala Lys Ala Glu Met Gln Leu Met Ser Pro Leu Gln Ile Ser Asp Pro Phe Gly Ser Phe Pro His Ser Pro Thr Met Asp Asn Tyr 25 Pro Lys Leu Glu Glu Met Met Leu Leu Ser Asn Gly Ala Pro Gln Phe 40 Leu Gly Ala Ala Gly Ala Pro Glu Gly Ser Gly Ser Asn Ser Ser Ser Ser Ser Ser Gly Gly Gly Gly Gly Gly Gly Gly Ser Asn Ser Ser Ser Ser Ser Ser Thr Phe Asn Pro Gln Ala Asp Thr Gly Glu Gln Pro Tyr Glu His Leu Thr Ala Glu Ser Phe Pro Asp Ile Ser Leu Asn Asn 105 Glu Lys Val Leu Val Glu Thr Ser Tyr Pro Ser Gln Thr Thr Arg Leu 120 Pro Pro Ile Thr Tyr Thr Gly Arg Phe Ser Leu Glu Pro Ala Pro Asn 135 130 Ser Gly Asn Thr Leu Trp Pro Glu Pro Leu Phe Ser Leu Val Ser Gly 155 Leu Val Ser Met Thr Asn Pro Pro Ala Ser Ser Ser Ser Ala Pro Ser Pro Ala Ala Ser Ser Ala Ser Ala Ser Gln Ser Pro Pro Leu Ser Cys 185 180 Ala Val Pro Ser Asn Asp Ser Ser Pro Ile Tyr Ser Ala Ala Pro Thr Phe Pro Thr Pro Asn Thr Asp Ile Phe Pro Glu Pro Gln Ser Gln Ala 210 Phe Pro Gly Ser Ala Gly Thr Ala Leu Gln Tyr Pro Pro Pro Ala Tyr 230 235 Pro Ala Ala Lys Gly Gly Phe Gln Val Pro Met Ile Pro Asp Tyr Leu Phe Pro Gln Gln Gly Asp Leu Gly Leu Gly Thr Pro Asp Gln Lys 265 Pro Phe Gln Gly Leu Glu Ser Arg Thr Gln Gln Pro Ser Leu Thr Pro

		275					280					285			
Leu	Ser 290	Thr	Ile	Lys	Ala	Phe 295	Ala	Thr	Gln	Ser	Gly 300	Ser	Gln	Asp	Leu
Lys 305	Ala	Leu	Asn	Thr	Ser 310	Tyr	Gln	Ser	Gln	Leu 315	Ile	Lys	Pro	Ser	Arg 320
Met	Arg	Lys	Tyr	Pro 325	Asn	Arg	Pro	Ser	Lys 330	Thr	Pro	Pro	His	Glu 335	Arg
Pro	Tyr	Ala	Cys 340	Pro	Val	Glu	Ser	Cys 345	Asp	Arg	Arg	Phe	Ser 350	Arg	Ser
Asp	Glu	Leu 355	Thr	Arg	His	Ile	Arg 360	Ile	His	Thr	Gly	Gln 365	Lys	Pro	Phe
Gln	Cys 370	Arg	Ile	Cys	Met	Arg 375	Asn	Phe	Ser	Arg	Ser 380	Asp	His	Leu	Thr
Thr 385	His	Ile	Arg	Thr	His 390	Thr	Gly	Glu	Lys	Pro 395	Phe	Ala	Cys	Asp	Ile 400
Cys	Gly	Arg	Lys	Phe 405	Ala	Arg	Ser	Asp	Glu 410	Arg	Lys	Arg	His	Thr 415	Lys
Ile	His	Leu	Arg 420	Gln	Lys	Asp	Lys	Lys 425	Ala	Asp	Lys	Ser	Val 430	Val	Ala
Ser	Ser	Ala 435	Thr	Ser	Ser	Leu	Ser 440	Ser	Tyr	Pro	Ser	Pro 445	Val	Ala	Thr
Ser	Tyr 450	Pro	Ser	Pro	Val	Thr 455	Thr	Ser	Tyr	Pro	Ser 460	Pro	Ala	Thr	Thr
Ser 465	Tyr	Pro	Ser	Pro	Val 470	Pro	Thr	Ser	Phe	Ser 475	Ser	Pro	Gly	Ser	Ser 480
Thr	Tyr	Pro	Ser	Pro 485	Val	His	Ser	Gly	Phe 490	Pro	Ser	Pro	Ser	Val 495	Ala
Thr	Thr	Tyr	Ser 500	Ser	Val	Pro	Pro	Ala 505	Phe	Pro	Ala	Gln	Val 510	Ser	Ser
Phe	Pro	Ser 515	Ser	Ala	Val	Thr	Asn 520	Ser	Phe	Ser	Ala	Ser 525	Thr	Gly	Leu
Ser	Asp 530	Met	Thr	Ala	Thr	Phe 535	Ser	Pro	Arg	Thr	Ile 540	Glu	Ile	Cys	

<210> 338 <211> 148 <212> PRT

<213> Homo sapiens

<400> 338

Pro Pro Ala Thr Ser Tyr Ala Pro Ser Asp Val Pro Ser Gly Val Ala
5 10 15

Leu Phe Leu Thr Ile Pro Phe Ala Phe Phe Leu Pro Glu Leu Ile Phe 20 25 30

Gly Phe Leu Val Trp Thr Met Val Ala Ala Thr His Ile Val Tyr Pro 35 40 45

Leu Leu Gln Gly Trp Val Met Tyr Val Ser Leu Thr Ser Phe Leu Ile 50 55 60

Ser Leu Met Phe Leu Leu Ser Tyr Leu Phe Gly Phe Tyr Lys Arg Phe 65 70 75 80

Glu Ser Trp Arg Val Leu Asp Ser Leu Tyr His Gly Thr Thr Gly Ile 85 90 95

Leu Tyr Met Ser Ala Ala Val Leu Gln Val His Ala Thr Ile Val Ser 100 105 110

Glu Lys Leu Leu Asp Pro Arg Ile Tyr Tyr Ile Asn Ser Ala Ala Ser 115 120 125

Phe Phe Ala Phe Ile Ala Thr Leu Leu Tyr Ile Leu His Ala Phe Ser 130 135 140

Ile Tyr Tyr His 145

<210> 339

<211> 196

<212> PRT

<213> Homo sapiens

<400> 339

Met Pro Gly Met Phe Phe Ser Ala Asn Pro Lys Glu Leu Lys Gly Thr
5 10 15

Thr His Ser Leu Leu Asp Asp Lys Met Gln Lys Arg Arg Pro Lys Thr 20 25 30

Phe Gly Met Asp Met Lys Ala Tyr Leu Arg Ser Met Ile Pro His Leu 35 40 45

Glu Ser Gly Met Lys Ser Ser Lys Ser Lys Asp Val Leu Ser Ala Ala 50 55 60

Glu Val Met Gln Trp Ser Gln Ser Leu Glu Lys Leu Leu Ala Asn Gln 65 70 75 80

Thr Gly Gln Asn Val Phe Gly Ser Phe Leu Lys Ser Glu Phe Ser Glu 85 90 95

Glu Asn Ile Glu Phe Trp Leu Ala Cys Glu Asp Tyr Lys Lys Thr Glu 100 105 110

Ser Asp Leu Leu Pro Cys Lys Ala Glu Glu Ile Tyr Lys Ala Phe Val 115 120 125

His Ser Asp Ala Ala Lys Gln Ile Asn Ile Asp Phe Arg Thr Arg Glu 130 135 140

Ser Thr Ala Lys Lys Ile Lys Ala Pro Thr Pro Thr Cys Phe Asp Glu 145 150 155 160

Ala Gln Lys Val Ile Tyr Thr Leu Met Glu Lys Asp Ser Tyr Pro Arg 165 170 175

Phe Leu Lys Ser Asp Ile Tyr Leu Asn Leu Leu Asn Asp Leu Gln Ala 180 185 190

Asn Ser Leu Lys 195

<210> 340

<211> 316

<212> PRT

<213> Homo sapiens

<400> 340

Met Ala Thr Phe Val Glu Leu Ser Thr Lys Ala Lys Met Pro Ile Val 5 10 15

Gly Leu Gly Thr Trp Lys Ser Pro Leu Gly Lys Val Lys Glu Ala Val 20 25 30

Lys Val Ala Ile Asp Ala Gly Tyr Arg His Ile Asp Cys Ala Tyr Val 35 40 45

Tyr Gln Asn Glu His Glu Val Gly Glu Ala Ile Gln Glu Lys Ile Gln 50 55 60

Glu Lys Ala Val Lys Arg Glu Asp Leu Phe Ile Val Ser Lys Leu Trp 65 70 75 80

Pro Thr Phe Phe Glu Arg Pro Leu Val Arg Lys Ala Phe Glu Lys Thr 85 90 95

Leu Lys Asp Leu Lys Leu Ser Tyr Leu Asp Val Tyr Leu Ile His Trp
100 105 110

Pro Gln Gly Phe Lys Ser Gly Asp Asp Leu Phe Pro Lys Asp Asp Lys

		115					120					125					
Gly	Asn 130	Ala	Ile	Gly	Gly	Lys 135	Ala	Thr	Phe	Leu	Asp 140	Ala	Trp	Glu	Ala		
Met 145	Glu	Glu	Leu	Val	Asp 150	Glu	Gly	Leu	Val	Lys 155	Ala	Leu	Gly	Val	Ser 160		
Asn	Phe	Ser	His	Phe 165	Gln	Ile	Glu	Lys	Leu 170	Leu	Asn	Lys	Pro	Gly 175	Leu		
Lys	Tyr	Lys	Pro 180	Val	Thr	Asn	Gln	Val 185	Glu	Cys	His	Pro	Tyr 190	Leu	Thr		
Gln	Glu	Lys 195	Leu	Ile	Gln	Tyr	Cys 200	His	Ser	Lys	Gly	Ile 205	Thr	Val	Thr		
Ala	Tyr 210	Ser	Pro	Leu	Gly	Ser 215	Pro	Asp	Arg	Pro	Trp 220	Ala	Lys	Pro	Glu		
Asp 225		Ser	Leu	Leu	Glu 230	Asp	Pro	Lys	Ile	Lys 235	Glu	Ile	Ala	Ala	Lys 240		
His	Lys	Lys	Thr	Ala 245	Ala	Gln	Val	Leu	Ile 250	Arg	Phe	His	Ile	Gln 255	Arg		
Asn	Val	Ile	Val 260	Ile	Pro	Lys	Ser	Val 265	Thr	Pro	Ala	Arg	Ile 270	Val	Glu		
Asn	Ile	Gln 275	Val	Phe	Asp	Phe	Lys 280	Leu	Ser	Asp	Glu	Glu 285	Met	Ala	Thr		
Ile	Leu 290	Ser	Phe	Asn	Arg	Asn 295	Trp	Arg	Ala	Cys	Asn 300	Val	Leu	Gln	Ser		
Ser 305		Leu	Glu	Asp	Tyr 310	Pro	Phe	Asn	Ala	Glu 315	Tyr						
<21	0> 3	41 <211 <212 <213	> DN.	A	apie	n											
		<220 <221 <222 <223	> mi > (1	)	(422	)											
ga	tgan	<400 attn	ttn	cnag	aga 🤉	gagg	aaga	ng c	tatt	cagt	t gg	atgg	gatt	aaa	tgcatca		60
ca	aata	aqaq	aac	ttag	aga 🤄	gaag	tcgg	aa a	agtt	tgcc	t tc	caag	cccg	aag	ttaacag gagaact		L20 L80
a a	itgat	gaaa aaat	gct	acca tctq	atq	cttt	artg agat	aa q	aaaa ataa	ggct	a ya a at	atca	ctga	ctg	atgaaaa		240
tg	ctct	ttct	gga	aatg	agg .	aact	aaca	gt c	aaaa	ttaa	g tg	tgat	aagg	aga	agacctg	3	300

ctgcatgtca cagacaccgg tgtaggaatg accagagaag agttggttaa aaaccttggt accatagcca aatctgggac aagcgagttt ttaaacaaaa tgactgaagc acaggaagat gg	360 420 422
<210> 342 <211> 472 <212> DNA <213> Homo sapien	
<220> <221> misc_feature <222> (1)(472) <223> n = A,T,C or G	
ctggagaagg tgtgcagggg aaaccctgct gatgtcaccg aggccaggtt gtcttctac tcgggacact cttcctttgg gatgtactgc atggtgtct tggcgctgna tgtgcaggca cgactctgtt ggaagtgggc acggctgctg cgacccacag tccagttctt cctggtggcc tttgccctct acgtgggcta cacccgcgtg tctgattaca aacaccactg gagcgatgtc cttgttggcc tcctgcaggg ggcactggtg gctgccctca ctgtctgcta catctcagac tcctcaaag cccgacccc acagcactgt ctgaaggagg aggagctgga acggaagccc agcctgtcac tgacgttgac cctggggag gctgaccaca accactatgg atacccgcac tcctcctct gaggccggac cccgccagg cagggagcta ctgtgagtcc ag	60 120 180 240 300 360 420 472
<210> 343 <211> 139 <212> DNA <213> Homo sapien	
<400> 343 gtcctgggcc ttccccttcc ctcaagccag ggctcctcct cctgtcgtgg gctcattgtg accactggcc tctctacagc acggcctgtg gcctgttcaa ggcagaacca cgacccttga ctcccgggtg gggaggtgg	60 120 139
<210> 344 <211> 235 <212> DNA <213> Homo sapien	
<pre>&lt;400&gt; 344 ctgcgggctc agcacagtag acatgactgg gatccccacc ttggacaacc tccagaaggg agtccaattt gctctcaagt accagtcgct gggccagtgt gtttacgtgc attgtaaggc tgggcgctcc aggagtgcca ctatggtggc agcatacctg attcaggtgc acaaatggag tccagaggag gctgtaagag ccatcgccaa gatccggtca tacatccaca tcagg</pre>	60 120 180 235
<210> 345 <211> 458 <212> DNA <213> Homo sapien	
<400> 345 ctgtaaggtg ctattcagtc ctgtgaccct tattttggaa tgctcttcat tactgttgct ctgttttgtg acttcctggg aaaccgccta ctttggtgtg gtgtcacctt gagctgtgca cataggacac cagttttgac ttaacctaac aggcagtttt tatctctagc tttttcaagc	60 120 180

caggtattga gcagtttett ggecaatgge etgagaaace acetgteeet gteaaggggt gattttattg gttttaagtg gggaagtaat eceatgtaet tatttettaa ataeetagga agttettett ggtggeteet ettggeeete ecetetttet eceeeaacee aeeateetge aaggeaagga atggeetete eeteeacaga ggeaaegget geagagggag eaetgtgget gecateecag tteetettea aageeaaaca gacaegeg	240 300 360 420 458
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gaaacgggtg ttccgaataa tatcctggaa gttatcagga cacctatttt aa	actatagge 240 acttaatte 300
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                                                                         180
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caagcataat atagcaagga ctaaccecta tacettetge ataatgaatt aactagaaat	180
aactttgcaa ggggagccaa agctaagacc cccgaaacca gacgagctac ctaagaacag	240
ctaaaagagc acacccgtct atgtagcaaa atagtgggaa gatttatagg tagaggcgac	300
aaacctaccg agcctggtga tagctggttg tccaagatag aatcttagtt caactttaaa	360
tttgcccaca qaaccctcta aatccccttg taaatttaac tgttagtcca aagaggaaca	420
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totoataaco otoaacacoo actooctott agocaatatt gtgootattg coatactagt	180
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212 424	
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(21) 10.110 Sup 2011	
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<pre>&lt;213&gt; Homo sapien  &lt;220&gt;</pre>	60 120 168
<pre> &lt;210&gt; 503   &lt;211&gt; 173   &lt;212&gt; DNA   &lt;213&gt; Homo sapien  &lt;220&gt;   &lt;221&gt; misc_feature   &lt;222&gt; (1)(173)   &lt;223&gt; n = A,T,C or G</pre>	
<pre>&lt;400&gt; 503 cctttataat aaattaggca aaaggttcag tgcnnggcta tantggacaa catgaaactc cataaaaatg actggatagg gggactgctt gagacttttc tttttgggcat tactaacaga attcaaagaa attccaacca cgcttatttt tccaaattct actgaaatga gag  &lt;210&gt; 504 &lt;211&gt; 310 &lt;212&gt; DNA &lt;213&gt; Homo sapien</pre>	60 120 173
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tgtgagaact ccctcagtat tagaagagca tgagggaaac cgcctccata atccaatcac
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ctctgctgtt ttggtcactg gtgcctagaa tttggggatg tacattggtt ttgattcaca
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cccactattt tgctacatag acgggtgtgc tcttttagct gttcttaggt agctcgtctg
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ccttgtagta ttgattgctc aatctataaa gagctcagtt tacagcatga ctgttagtaa
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cagcccagta gtctaacatt aagggtctta ggaaatgaga acttatcacc tttcctt atgaaaaggt aacctccagg taaccaaaaa tagaacttcc tctgtgttcg ttttta	
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nttgttcctg anaanggtgg gtaaatattc caggaaaaan aatgcacatt gaatgg	atgt 300
gagagaccac attgcctctc ccactgcttt ggggagcact ttcctgtcat ttctaa	ctta 360
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tatttcctga gcgtctgaga tgttagtatt agttagtttt gttgtaagng ttaggaaaag
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atcttaaaaa ggaactttaa atgcagggct atattgaatt ggnaaactgc aacacaaact
ggcgcaacat aggtaaatga ataccaatct cactctatgt gatgcaagca tgctactttc
                                                                          180
ccactaattt aaattacttt caaccactat gagccagaat gcatgcctga accttaaact
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gcactttaaa aagtaacatc ttggcctaa
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      <211> 266
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                                                                          120
                                                                          180
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cgattagttt tagcattgga gtaggtttag gttatgnacc gtactctagg ccatatgtgt
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tgganattga nactagtagg gctagg
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agttgtagga gatgagatca aaggctagga atgaagtgta aggccatgtc atgtgacctt
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atatcccagg attttggtta ttcatgcctt tcttttgtga ctttctttca aattagccaa
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ttaaagatac cccttcaatc accggtgaca tcagtacaac agtttttcaa cagttttctc
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cattggacca ataggttaca ttttcgttcc ttttttgttt tggttcatct gttaagcagt
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cttggaatgc tattactgtt cacacaaagt atgattctgt ttgaataagg caaatgctcc
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                                                                        240
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                                                                        300
gggcttgatt ttttttaaaa aaacagaatg aattgatgtc ttattttata aatgttctat
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                                                                        180
tagaaggtaa agcttgagaa gatgagggtg tttacgtaga ccagaaccaa tttagaagaa
                                                                        240
tacttgaagc tagaagggga agttggttaa aaatcacatc aaaaagctac taaaaggact
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                                                                        120
ggatgtaatg ctattttaat cctgctgtgg cccccaatac tagtacttgt ccataccttc
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ttgcattttt agcgtctgct ctgtggggtt gttaggccct ggcactccca ggaactagtg
                                                                        240
ctaaagctgc atctntctct cccctctagg gatcgataaa gtttcactgc agaaagtctc
                                                                        300
                                                                        360
cactgoggta tgctgacato tgccctgaac cttcacccta cagcattaca ggctttaatc
                                                                        420
agattetget ggaaagacae aggetgatee aegtgaeete ttetgeette aetgggetgg
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aagagcatgt aacagagaag aaggactana cctaccattt tctggataaa gaattggaaa
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                                                                         240
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                                                                         480
                                                                         540
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cggaagcgat taagaaaagg gtgggatgga aaaattaacc caacaatatt agatcaatac
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gtagtattta agngtccata atgtgccagg ctgaagatgc acgggaaaac cacactagcc
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gatacacaac agttaagcgt aaagatcaca ggcaatagca ttcaaacatg gatgtgggta
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                                                                        240
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agctatcata aaattcactt tcctgaagac atttactctc attcacttcc aaactccaaa
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cctttttctg gtagcaccac ttttgttttt aatagaaaga tgagttcata tctgtacatc
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• • • • • • • • • • • • • • • • • • • •	
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                                                                        360
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atcaaatttt tggtgcttgg tgcagagacg gcgtggggaa gggtgaatgg ttttgggaat
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                                                                        621
aattcagtgc acacctgggg g
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aactgattaa tgtttgggnn tgagtttnta tatcacagcc anaattntat gatgnaccat
gtancgaaca atgctacagg gatgaatatt atggagaagt antctanttt gaagcttagg
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aaatgaggaa attattggta accaatttat tttaaaagcc catcaattta atttctggtg	180 240
gtgcagaagt tagaaggtaa agcttgagaa gatgagggtg tttacgtaga ccagaaccaa tttagaagaa tacttgaagc tagaagggga agttggttaa aaatcacatc aaaaagctac	300
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                                                                         180
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                                                                         300
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                                                                         360
                                                                         420
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agaatactgt ggtgtggttg acagtagtgc ctaagtctgt tttcagagtg aaaaatgaca
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      <223> n = A, T, C \text{ or } G
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                                                                          60
cttagaggtg gggagcagag aattctctta tccaacatca acatcttggt cagatttgaa
                                                                         120
ctcttcaatc tcttgcactc aaagcttgtt aagatagtta agcgtgcata agttaacttc
                                                                         180
caatttacat actctgctta gaatttgggg gaaaatttag aaatataatt gacaggatta
                                                                         240
ttggaaattt gttataatga atgaaacatt ttgtcatata agattcatat ttacttctta
                                                                         300
tacatttgat aaagnaaggc atggttgtgg ttaatctggt ttatttttgn tccacaagtt
                                                                         360
                                                                         391
aaataaatca taaaacttga acaaaaaaaa a
      <210> 698
      <211> 536
      <212> DNA
      <213> Homo sapien
      <220>
      <221> misc feature
      <222> (1)...(536)
      \langle 223 \rangle n = A,T,C or G
      <400> 698
ctgagcatac agcaataaaa ataacataat ttttatgtgt acaatattta tggaatacgt
                                                                          60
tactggaaca gataaataat ttagttaata acatgacaaa gaacagaaat tgtatacact
                                                                         120
atacagcata gtaatagaat aatgaatgat taaagttatt aatattaggt agaaaatgaa
                                                                         180
                                                                         240
gggtatcttt gagagcagaa ctcaaggaag caagcaattt gccttatgag gaaagagtta
cctgtggata aaggagaaac tgaaaaattt acaagtcaag actttttgag caaagacaaa
                                                                         300
aatatgacta tgagtcacca attcagtaca gtgaaaaaaa agttgaagag atatcttgga
                                                                         360
agtaaaccat gttgtggaag agcagggttt tgataatcat gggattattc tgaatgaatt
                                                                         420
ttaaatgcga taggaatata tgagataatt tcaccagaga ataatatgat catgtttgca
                                                                         480
tttcaaaggg gtgtatctgg tgcactgngt agaataaata ggntatgtga gcaagt
                                                                         536
      <210> 699
      <211> 419
      <212> DNA
      <213> Homo sapien
      <220>
      <221> misc_feature
      <222> (1)...(419)
```

## <223> n = A,T,C or G

attgagacta gtagggctag g

<223> n = A,T,C or G	
4005 600	
<400> 699 ngtccacctg agggcaggtg acaaggacct gacagagccc atgcagggct ttagatttgg	60
acacacaaga gttgataact teeteatgaa eteettgeet gatetaaact catattatgg	120
gttctgactg tttgagtaat catcttcaag gttaaacctc ttggcagtta cccttttcac	180
aaagtgcaca gtgggaatcg agaatcgata gggttaattt tggagcagtg gcttatacca	240
ttcacctctg tttttttgtg attatttcac agataatgag accttaataa caaataggcg	300
taaaaaaatt ttcacattga aatgatagaa acatttgatg taataaaact tggttggctt	360
gatattttaa ggaattgaaa cctagcaatc ttattggaga gacaagaatt ggtctccag	419
<210> 700	
<211> 336	
<212> DNA	
<213> Homo sapien	
<400> 700	60
ccacttattg tccttaaaaa tccatactga tacatggaca gtaagtgtgt tttcagatgg	120
agtaccagca ccgaaaatgg gttgagggag gatgggttgt atgtatgttt ctgcccacta attttgagca gccatattat gaattaaatc gtcacagcca agtaataacc caagaatggt	180
attetgagea gecatattat gaattaaatt gecatageta ageateadee edagadegge atgagettea tgtgtaatag etcaaatgga ataageatga atgetggagt ggaccattat	240
cctcaaatat tctatgtcac ttctcattta aagactcttg ttatgaacta ttagaaactt	300
taggcaaaat caaaagtatt tgcggcaaaa taaagg	336
cayycaaaat caaaaycacc cycyyoanaa cannoo	
<210> 701	
<211> 418	
<212> DNA	
<213> Homo sapien	
<400> 701	60
ccatgtgatg atgttgacaa cccctgaaga gcctcagtcc attgttccac gtttaagaac	60 120
taggaatacc aggactgatg caattctact gggtcactat cgcttgtcac aagacacaga	180
caatcagacc aaagtatttg ctgtaataac taagaaaaaa gaagaaaaac cacttgacta	240
taaatacaga tattttegte gtgteeetgt acaagaagea gateagagtt tteatgtggg getacageta tgtteeagtg gteaceagag gtteaacaaa eteatetgga tacateatte	300
ttgtcacatt acttacaaat caactggtga gactgcagtc agtgcttttg agattgacaa	360
gatgtacacc cccttgttct tcgccagagt aaggagctac acagctttct cagaaagg	418
gatgtatate teetigetee tegeouguge unggingerie nengere	
<210> 702	
<211> 261	
<212> DNA	
<213> Homo sapien	
<220>	
<221> misc_feature	
<222> (1)(261)	
<223> n = A, T, C  or  G	
400 702	
<400> 702 gggcctgttg tgggggtggg ggaagcaggg aggggaacag ctaaataggt tgctgttgat	60
ttggttaaaa aatagtaggg ggatgatgct aataattagg ctgngggtgg ttgtgttgat	120
tcaaattatg tgttttttgg agagtcatgt cagtggtaga aatataattg ttgggacnat	180
tagntttagc attggagtag gtttaggtta tgtacgtagt ctaggccata tgtgttggan	240

<211> 266

```
<210> 703
      <211> 261
      <212> DNA
      <213> Homo sapien
      <220>
      <221> misc_feature
      <222> (1)...(261)
      <223> n = A, T, C or G
      <400> 703
gggcctgttg tgggggtggg ggaagcaggg aggggaacan ctaaataggt tgctgttgat
                                                                        60
ttggttaaaa aatagtaggg ggatgatgct aataattagg ctgngggtgg ttgtgttgat
                                                                        120
                                                                        180
tcaaattatg tgttttttgg agagtcatgt cagtggtagt aatataattg ttgggacnat
tagntttagc attggagtag gtttaggtta tgtacgtagn ctaggccata tgtgttggag
                                                                        240
                                                                        261
attganacta gtagggctag g
      <210> 704
      <211> 381
      <212> DNA
      <213> Homo sapien
      <220>
      <221> misc_feature
      <222> (1)...(381)
      <223> n = A,T,C or G
      <400> 704
ngtntgaatt ctattaaaga tacaaagagg agctggtacc atttcttctg aaactattac
                                                                         60
aaacaactga aaaggtggaa tttctcccta attcatttta ggaggccagc attatactga
                                                                        120
                                                                        180
taccaaaacc tggcagaggt acaataataa aaggaaactt caagtcagta tcactgatga
acaccaatgt gaaaatcctc aataaaatac tggcaaactg aattcagcag cacatcaaaa
                                                                        240
agctaatcca ccacaatcaa gtcagcttca tccctgcgat gcaagtctgg ttcaacatat
                                                                        300
gcaaatcaat aaatacaatt catcagataa acagagctaa agacaaaatt cacatgattt
                                                                        360
                                                                        381
tctcaataga tgcagaaaag g
      <210> 705
      <211> 477
      <212> DNA
      <213> Homo sapien
      <400> 705
ctgaaccctc gtggagccat tcatacaggt ccctaattaa ggaacaagtg attatgctac
                                                                         60
ctttgcacgg ttagggtacc gcggccgtta aacatgtgtc actgggcagg cggtgcctct
                                                                        120
aatactggtg atgctagagg tgatgttttt ggtaaacagg cggggtaaga tttgccgagt
                                                                        180
tccttttact ttttttaacc tttccttatg agcatgcctg tgttgggttg acagtgaggg
                                                                        240
taataatgac ttgttggtga ttgtagatat tgggctgtta attgtcagtt cagtgtttta
                                                                        300
                                                                        360
atctgacgca ggcttatgcg gaggagaatg ttttcatgtt acttatacta acattagttc
ttctataggg tgatagattg gtccaattgg gtgtgaggag ttcagttata tgtttgggat
                                                                        420
tttttaggta gtgggtgttg agcttgaacg ctttcttaat tggtggctgc ttttagg
                                                                        477
      <210> 706
```

```
<212> DNA
     <213> Homo sapien
     <220>
     <221> misc_feature
     <222> (1)...(266)
     <223> n = A, T, C \text{ or } G
      <400> 706
60
ggaggttagt tgtggcaata aaaatgatta aggatactan tataagagat caggntcgtc
                                                                      120
ctttagtgtt gtgtatggct atcatttgtt ttgaggntag tttgattagt cattgttggg
                                                                      180
                                                                      240
tggtaattag teggttgttg atgagatatt tggaggtggg gateaataga gggggaaata
                                                                      266
gaatgatcag tactgcggcg ggtagg
      <210> 707
      <211> 358
      <212> DNA
      <213> Homo sapien
      <220>
      <221> misc_feature
      <222> (1)...(358)
      \langle 223 \rangle n = A,T,C or G
      <400> 707
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                                                                       60
caatcattaa aaagtcagga aacaacaggt gctggagagg atgtggagaa ataggaacac
                                                                      120
ttttacaccg ntggtgggac tgtaaactag ttcaaccatt gtggaagtca gtgtggcgat
                                                                      180
tecteaagga tetagaacta gaaataccat ttgacccage eggecaatat teaacattet
                                                                      240
taaaggaaag aattttcaac ccagaatttc atatccagcc aaactaagct tcgttagtga
                                                                      300
aggagaaata aaatacttta cagacaagca aatactgaga gattttgtca ccaccagg
                                                                      358
      <210> 708
      <211> 491
      <212> DNA
      <213> Homo sapien
      <220>
      <221> misc feature
      <222> (1)...(491)
      <223> n = A, T, C \text{ or } G
      <400> 708
cctactatgg gngttaaatt ttttactctc tctacaaggt tttttcctag tgtccaaaga
                                                                       60
                                                                       120
gctgttcctc tttggactaa cagttaaatt tacaagggga tttagagggt tctgtgggca
                                                                       180
aatttaaaqt tgaactaaga ttctatcttg gacaaccagc tatcaccagg ctcggtaggt
ttgtcgcctc tacctataaa tcttcccact attttgctac atagacgggt gtgctctttt
                                                                       240
                                                                       300
agctgttctt aggtagctcg tctggtttcg ggggtcttag ctttggctct ccttgcaaag
                                                                       360
ttatttctag ttaattcatt atgcagaagg tataggggtt agtccttgct atattatgct
tggttataat ttttcatctt tcccttgcgg tactatatct attgcgccag gtttcaattt
                                                                       420
ctatcgccta tactttattt gggtaaatgg tttggctaag gttgtctggt agtaagggng
                                                                       480
                                                                       491
gagtgggttt g
```

```
<210> 709
      <211> 460
      <212> DNA
      <213> Homo sapien
      <220>
      <221> misc feature
      <222> (1)...(460)
      <223> n = A, T, C or G
      <400> 709
nggttttttt tgtagagcaa ataatttatg caaaatatgt tacaaaatct gggatgctaa
                                                                         60
atagttgaca caagtactgt gtttgacatt tagtttcatt tgaattagta atagaatttg
                                                                         120
ctccttccaa catttacatc ttttttcttt ctgactttat atattttcaa taaaaatttg
                                                                         180
ctccacagtt tttaagntca ttcttcttga atccgntttt acatttgctg ngacaaacct
                                                                         240
gcataaaact agattttata gatataactt ctttggaaga gataaaaatt caaaagtttg
                                                                         300
acattgettt canttattet tttetteatt gttttgattg geceetgtta gattgatgta
                                                                         360
ttgccaatct acttttgatg gcatgaatnt aaaatgacaa cataaaaagc ncttctagtg
                                                                         420
caacagtaat tgaaacttgc agttttccat taaaaaaaaa
                                                                         460
      <210> 710
      <211> 542
      <212> DNA
      <213> Homo sapien
      <220>
      <221> misc_feature
      <222> (1)...(542)
      \langle 223 \rangle n = A,T,C or G
      <400> 710
ctgttacagt gacaagagat aaaaagatag acctgcagaa aaaacaaact caaagaaatg
                                                                          60
                                                                         120
tgttcagatg taatgtaatt ggagtgaaaa actgtgggaa aagtggagtt cttcaggctc
ttcttggaag aaacttaatg aggcagaaga aaattcgtga agatcataga tcctactatg
                                                                         180
cgattaacac tgtttatgta tatggacaag agaaatactt gttgttgcat gatatctcag
                                                                         240
                                                                         300
aatcggaatt tctaactgaa gctgaaatca tttgngatgt tgtatgcctg gtatataatg
tcagcaatcc caaatccttt gaatactgtg ccaggatttt taagcaacac tttatggaca
                                                                         360
                                                                         420
gcagaatacc ttgcttaatc gtagctgcaa agtcagacct gcatgaagtt aaacaagaat
acagtatttc acctactgat ttctgcagga aacacaaaat gcctccacca caagccttca
                                                                         480
cttgcaatac tgctgatgcc cccagtnagg atatctttgt taaattgaca acaatggacc
                                                                         540
                                                                         542
tg
      <210> 711
      <211> 394
      <212> DNA
      <213> Homo sapien
      <220>
      <221> misc feature
      <222> (1)...(394)
      <223> n = A, T, C \text{ or } G
      <400> 711
caaacccact ccaccttact accagacaac cttagccaaa ccatttaccc aaataaagta
                                                                          60
```

taggcgatag aaattgaaac ctggcgcaat agatatagta ccgcaaggga aagatgaaaa	120
attataacca aqcataatat agcaaggact aacccctata ccttctgcat aatgaattaa	180
ctanaaataa ctttgcaagg agagccaaag ctaagacccc cgaaaccaga cgagctacct	240
aagaacaqct aaaagagcac acccgtctat gtagcaaaat agtgggaaga tttataggna	300
gaggegacaa acctacegag cetggtgata getggttgte caagatagaa tettagttea	360
actttaaatt tgcccacaga accctctaaa tccc	394
<210> 712	
<211> 552	
<212> DNA	
<213> Homo sapien	
<220>	
<221> misc_feature	
<222> (1) (552)	
$\langle 223 \rangle$ n = A,T,C or G	
400 710	
<400> 712 gaggtctgta naatgccagg ctcaaatttg tctttataat ttaataccag aaatctttcc	60
cttgtgatgt ttetttettt etggattgee tetatageag gggatagegg gggaggataa	120
qqcacatctt tgntgtactg agaaatttga ccacgcagga tgatgtggct gttctcattc	180
atctgcacag agaaaaataa tgataaaata tccctttcct atgtttactg attttatggc	240
tgccataatg gaagcctcct tgactattta atcctttctg tcaactaggt tcgattttt	300
ttttaattta cetgttagag gtatttaana attttaacta getanaaata attacattee	360
aaaggaacac caaggcaaat aaatggttgg taatcagcaa aagaattaca ttagttgttg	420
ntgctactta ttagggggag aactgttttt ttttaaattt aaacaattta ataatctcaa	480
ctgcaaataa ttttagatgc agcaaaggac tatgtagncg ttaatacctc atgttgatat	540
_	552
tttcataata tt	332
<210> 713	
<211> 518	
<212> DNA	
<213> Homo sapien	
(213) Nome Suppose	
<220>	
<221> misc_feature	
<222> (1)(518)	
<223> n = A,T,C or G	
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ccaaaaactg gaagcagctc actaaacaaa cagtggcata cccatagaac tgcatacttc	120
tcagcagtat gaaagaatga gctacttata taagcatcat tgataaacct caaaaaaaaa	180
atgccacatg aanaaaccca aagggganaa acataaaaac tttatatgtc agtcatataa	240
aattetanaa aatgeaaact aateeatent aaaggaaagt aaateaacag ttgtetggag	300
gaccananag agcaggagga ganagattat taaaggggtt aaagtaaatt tgggagtgcc	360
cttccntttt taaatnctat gaaaatgaaa gtaaaggcnc atgcatgttg taaactaata	420
gtaacaaaca naatgggttg gagtggggtg ttgtctgggg acatcattac aaaatgtaag	420
ccagtttatn taaattttga aaagaccgtg gactctgatc tgactgatna atgttggaag	480 518
agataagtgt gctgcaaatg ggggaattaa taaaacag	эта
<210> 714	
<211> 281	
<211> 281 <212> DNA	
CZIZZ DIM	

<213> Homo sapien

<pre>&lt;400&gt; 714 ccaattgatt tgatggtaag ggagggatcg ttgacctcgt ctgttatgta aaggatgcgt agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct atttcctgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg gcatacagga ctaggaagca gataaggaaa atgactatga gggcgtgatc atgaaaggtg ataagctctt ctatgatagg ggaagtagcg tcttgtagac c  &lt;210&gt; 715 &lt;211&gt; 443 &lt;212&gt; DNA</pre>	60 120 180 240 281
<213> Homo sapien <400> 715 cttgaaatca gcaacacact tacaaatgag aaaatgaaaa tagaagagta tataaagaaa gggaaagagg attatgaaga gagtcatcag agagctgtgg ctgcagaggt atccgtactt gaaaactgga aggagagtga agtgtataag ctacagatca tggagtcaca agcagaagcc tttctgaaga agctggggct gattagccgt gatcctgcag catatcccga catggagtct gatatacgtt catgggaatt gttctttct aatgttacaa aagaaattga gaaagcaaag tctcagtttg aagaacaaat taaggcaatt aaaaatggtt cccggctcag tgaactttct aatgtgagtct catgggtctt caggccacga tgg	60 120 180 240 300 360 420 443
<210> 716 <211> 639 <212> DNA <213> Homo sapien  <220> <221> misc_feature <222> (1)(639) <223> n = A,T,C or G	
<pre>&lt;400&gt; 716  ccaaanaaaa tgaagtacag agtctgcata gtaagcttac agataccttg gtatcaaaac aacagttgga gcaaagacta atgcagttaa tggaatcaga gcagaaaagg gtgaacaaag aagagtctct acaaatgcag gttcaggata ttttggagca gaatgaggct ttgaaaggtc aaattcagca gttccattcc cagatagcag cccagacctc cgcttcagtt ctagcagaag aattacataa agtgattgca gaaaaggata agcagataaa acagactgaa gattctttag caagtgaacg tgatcgtta acaagtaaag aagaggaact taaggatata cagaatatga attcttatt aaaagctgaa gtgcagaaat tacaggccct ggcaaatgag caggctgctg ctgcacatga attggagaag atgcaacaaa gtgtttatgt taaagatgat aaaataagat tgctggaaga gcaactacaa catgaaattt caaacnaaat ggaagaattt angattctaa atgaccaaaa canagcatta aaatcagaag gaaaaattg gcagactctt gtttctgcac angcctaata aggatgntgn ggaacaaatg gaaaaattg</pre>	60 120 180 240 300 360 420 480 540 600 639
<210> 717 <211> 473 <212> DNA <213> Homo sapien  <220> <221> misc_feature <222> (1)(473)	

## $\langle 223 \rangle$ n = A,T,C or G <400> 717 nntgaggeta etgetgtttt attacaacat tacetettgt ttttataaag tgtaccaaga 60 tttaaattga taactttatt ttacttgaaa aaaaaaagtt tnttttatca ccagtgttac 120 agttgtcttc tgtttctttt tgttttgntt tatttgnttt cctttttagc caaagagtga 180 acagaanatt ttcttatttt ggtggctatt cattttactt ttaaaagtga ttggtggatt 240 300 ttagactaat tatgggggaa tttgccacca aaataaaaaa tatgtaaagn gtagtgatta cagagtggtt aaaatgtggg ttagtactta tttattccat taattgatta tttgactgtt 360 tataaagaaa gttgctttat ttctttaaac atcttcaaaa gatgatcctt tcttgtcaca 420 ttatagccaa aagaagcaga gaacttcact gtctgcattt ggttcctggt tgg 473 <210> 718 <211> 207 <212> DNA <213> Homo sapien <400> 718 ggtaaatgct agtataatat ttaccatctc acttctagga atactagtat atcgctcaca 60 cctcatatcc tccctactat gcctagaagg aataatacta tcactgttca ttatagctac 120 180 teteataace eteaacacee actecetett agecaatatt gtgeetattg ecatactagt 207 ctttgccgcc tgcgaagcag cggtagg <210> 719 <211> 255 <212> DNA <213> Homo sapien <220> <221> misc feature <222> (1)...(255) <223> n = A,T,C or G <400> 719 60 cctatattac ggatcatttc tctactcaga aacctgaaac atcggcatta tcctcctgct tgcaactata gcaacagcct tcataggcta tgtcctcccg tgaggccaaa tatcattctg 120 aggggccaca gtaattacaa acttactatc cgccatccca tacattggga cagacctagt 180 240 tcaatgaatc tgaggaggct actcagtaga cagncccacc ctcacacgat tctttacctt 255 tcacttcatc ttgcc <210> 720 <211> 455 <212> DNA <213> Homo sapien <220> <221> misc_feature <222> (1)...(455) <223> n = A, T, C or G<400> 720 ccaatgtcga aacctacaag atttccttaa aatctctaat agaggcatta cttgctttca 60

attgacaaat gatgccctct gactagtaga tttctatgat ccttttttgt cattttatga atatcattga ttttataatt ggtgctattt gaanaaaaaa atgtacattt attcatagat 120

180

agataagtat caggtctgac cccagtggaa aacaaagcca aacaaaactg aaccacaaaa aaaaaggctg gtgttcacca aaaccaaact tgttcattta gataatttga aaaagctcca tagaaaaggc gtgcagtact aagggaacaa tccatgtgat taatgnttnc attatgttca tgtaanaagc cccttatttt tagccataat tttgcatact gaaaatccaa taatcagaaa agtaattttg ccacattatt tatnaaaaat gttcc	240 300 360 420 455
<210> 721 <211> 530 <212> DNA <213> Homo sapien	
<220> <221> misc_feature <222> (1)(530) <223> n = A,T,C or G	
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<210> 722 <211> 242 <212> DNA <213> Homo sapien  <220> <221> misc_feature	
<222> (1)(242) <223> n = A,T,C or G	
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attetgtaac tacatacett tgaaacacta tteacattea aataaacget tgtttetag ceaggeacag geteaattag tttteaaac tetageeaag geagtatte atttgggaaa teatgeaaca gaactgetea attettaact teteetgetg ttaacattta eacttagaet geageaaca gttaacttaa attttggtet eaagggaaca aaaaaaaatt geatteagaa ttaatatag tatttaaaa etaatttag eetgtaagne attatgagea atagtaactt teatacetee teatettgne tgataatata ttetatatge tgneaatetg attatatagt etatatgeta gaagttgetg attteatte tgeeaceaaa aaaactgte ettttttt tatgggggaa aaagggaatt taaa	180 240 300 360 420 480 540
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atcaaqaqta attaccaact taatgttttt gcattggact ttgagttaag attattttt
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aaatcctqaq gactagcatt aattgacgg
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                                                                        120
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                                                                        180
qatattttta atatactggg ttacataaac tgttattaag attaattttg cttgtttctt
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ttttaagcac tctggggtac taacttgaac tcccagaaac ccataagcac actctgcata
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aattcacaaa atattgantc cttaacaaat gtcaattagt atatggagag agctaaagga
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ggagtttccg atgccagagg atgaaagcaa gtgctttctc caccctctcc		180
aaaacaaatc cttttgctga tacttgtttc aaaagcatcc attgtaaagc		240
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tcagtgctct gaattcaact gacagactaa agggtgtttc ctgtaacagt		360
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                                                                        240
cactacactt tacatatttt ttatttnggn ggcaaantcc cccataatta gtctaaaatc
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caccaatcac ttttaaaagt aaaatgaata gccaccaaaa taagaaaatc ttctgttcac
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                                                                        420
cactggtgat aaaagaaact ttttttttac aagtaaaata aagttatcaa tttaaatctt
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                                                                        300
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atttgcatct gggaaataga gaaagtacat ataagccatg attttgaagn caaaagagag
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agantatttg ccaaggaggg gtgagttata gtatgtaatt ataacataca gaagcttttt
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ggccacctaa tttgaaatca catatagacc aggcgcggtg gctcacgcct gtaattccaa

cactttggaa ggccaaggca ggtggatcac aaggtcaaga gattgagacc atcttggcca acatggcgaa accccgtctc taccaaaaat acaaaaatca g	300 341
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gaageettte tgaagaaget ggggetgatt ageegtgate etgeageata teeegacatg gagtetgata taegtteatg ggaattgttt etttetaatg ttacaaaaga aattgagaaa geaaagtete agtttgaaga acaaattaag geaattaaaa atggtteeeg geteagtgaa etttetaaag ngeagatte tgagetttea ttteetgeet gtaacaeggt teateeegag ttaeteeetg agtetteagg ceaegatgg	240 300 360 420 449
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ttgaaccagn ccagtctgat tttcaggtga gaagacagaa ggaattaggg aaaagggtga gataatggta tttttggtag ccacagggaa cgcacacgca cacacacaaa cacacaca	tacttacaga atagcaggag cgctaaaact aagtagatac cttcaaatgt	gtaaaggaaa gggactggag caaactaaaa taagggcaag	taaatgaaaa atcacacaca acctcccaaa aatagaccag	240 300 360 420 480 540 574
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cataaaatct ttgggaaggg acaactgtaa aggaagttca tagtcgtcaa tatgaaggat
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tttaatttet ggettteeta tettettett caggataget teetteagea tagaattgtt
                                                                         180
ttccaatata aaatattttg ctgggttgtc cgtactatgt aggctgacca ctgggaccct
                                                                         240
tggaccttca cagaataata agaaatgttg attcatggga ctaaaactgg catcaaaata
                                                                         300
                                                                         360
tgtacattgt tettteatga aattacatga aatgeattgg egatteaata ateetteagt
                                                                         375
agaagcactg tacag
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      <211> 485
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                                                                          60
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                                                                         180
tgcctctaat actggtgatg ctagaggtga tgtttttggn aaacaggcgg ggtaagattt
                                                                         240
geogagities tittaettit tittaasetti settatgage atgestgigt tgggttgasa
gtgagggtaa taatgacttg ttggtgattg tagatattgg gctgttaatt gtcagttcag
                                                                         300
                                                                         360
tgttttaatc tgacgcaggc ttatgcggag gagaatgttt tcatgttact tatactaaca
                                                                         420
ttagttette tatagggtga tagatnggte caattgggtg tgaggagnte aettatatgt
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ttang
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                                                                         120
ccaaaacctg gcagaggtac aataataaaa ggaaacttca agtcagtatc actgatgaac
                                                                         180
                                                                         240
accaatgtga aaatcctcaa taaaatactg gcaaactgaa ttcagcagca catcaaaaag
                                                                         300
ctaatccacc acaatcaagt cagcttcatc cctgcgatgc aagtctggtt caacatatgc
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                                                                         379
tcaatagatg cagaaaagg
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                                                                         60
atatgggtag aatatagtca atatgaatgg aatagacaat gctttgaaaa tcactggagg
                                                                        120
                                                                        180
gaggetttat tgtttgtgaa aacatgttgt catcactttt tgetttaage cettggtggt
                                                                        240
gaaataactc aaaccattct tccttatgct gaagatcgag aaccccaagt atcacatcta
                                                                        300
ccatcccact catcaatgtg attggtcagt ctttgctgag gncctgcata gccagtttta
aagttagagt tettgeatat acatatgaaa aggeatgtta ettgtgettt caaagagett
                                                                        360
                                                                        420
tttgcttggt gtaaaaagaa aactcaaatt acagtgtgat gtggaatata atggtggtag
                                                                        480
tttcatcgag atgatgggaa agaattgata agataaagcn gaaagatgag cagaattttc
                                                                        518
agattgggtn tggaaagagc acttaagaaa gagggtgg
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                                                                         60
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agaataagag ttaagaagaa aaagactatg agaaaggaag tgntgacccc atttgcattt
                                                                        180
aaatggcagg aatagtetea atetaeteat tggggaaaaa tgtatgttge atatttttga
                                                                        240
                                                                        300
gatattgcaa cttgctctct ctctttgcca ccccaccctt tgncatgctc tgtttttggg
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cttcttcctt ctttctcc
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cctcatatcc tccctactat gcctagaagg aataatacta tcactgttca ttatagctac
                                                                        120
teteataace eteaacacee acteeetett ageeaatatt gtgeetattg eeataetagt
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ctttgccgcc tgcgaagcag cggtagg
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      <210> 772
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<pre></pre>	~220s	
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His Lys 290	Arg Thr	Pro Ala	Leu 295	Ile	Ala	Leu	Arg	Tyr 300	Gln	Leu	Gln	Arg
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Asn Val	Gln Val	Phe Glu 325	Phe	Gln	Leu	Thr 330	Ser	Glu	Glu	Met	Lys 335	Ala
Ile Asp	Gly Leu	Asn Arg	Asn	Val	Arg 345	Tyr	Leu	Thr	Leu	Asp 350	Ile	Phe
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Arg Ile Gln Leu Arg Ser Gln Val Leu Gly Arg Glu Met Arg Asp Met

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85

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Gly Arg Arg Val Pro Arg Arg Arg Arg Trp Gly Cys Val Gln 65 70 75 80

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Asp Cys Arg Pro Phe Leu Ala His Ser Ala Gly Tyr Ile Leu Gly Ser 50 55 60

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Phe Cys Ser Lys Thr Lys Ala Leu Ala Ala Ile Pro Pro Pro Val Pro 65 70 75 80

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Gly Ile Thr Ala Leu Leu Asn Val Ser Ser Asp Cys Pro Asn His Phe 130 135 140

Glu Gly His Tyr Gln Tyr Lys Cys Ile Pro Val Glu Asp Asn His Lys 145 150 155 160

Ala Asp Ile Ser Ser Trp Phe Met Glu Ala Ile Glu Tyr Ile Asp Ala 165 170 175

Val Lys Asp Cys Arg Gly Arg Val Leu Val His Cys Gln Ala Gly Ile 180 185 190

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<223> n=A,T,C or G
<400> 812
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gctgccgagg gccgcgcggt gtacgtggtg gacgacgcag ctgtcctggg cgcagaggac 180
ccageggtgt acggcgattc tgcccgtgag aaggcattgc gtggagctct gcgagcctcc 240
gtggaacggc gcctgagtcg ccacgacgtg gtcatcctgg actcgcttaa ctacatcaaa 300
ggtttccgtt acgageteta etgeetggea egggeggege geacceeget etgeetggte 360
tactgcgtac ggcccggcgg cccgatcgcg ggacctcagg tggcgggcgc gaacgagaac 420
cctggccgga acgtcagtgt gagttggcgg ccacgcgctg aggaggacgg gagagcccag 480
geggegggea geagegteet eagggaactg catactgegg actetgtagt aaatggaagt 540
gcccaggccg acgtacccaa ggaactggag cgagaagaat ccggggctgc ggag
```

<210> 813 <211> 561

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<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(561)
<223> n=A,T,C or G
<400> 813
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tagatgaagc caaacattgt tggaggtact gaaatettag aetecaceat gtgtecagga 120
neceattgae gteetetett etgaaaaete egtgtggeee tegetetgea etgteatgag 180
geggtgatgg agetagatae ceaceaegga caatgateat eagtttgggg ttetetgggt 240
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acgggatete teatecagge gatacgtetg gteetgtgge atgtggetet enacgaaaca 360
ccagggangc attatgttgg ggacttettg gggetetget ggtetetget ccagacacga 420
ttaatccgaa atgtgttaan tcgancacat gggtccacgt ccaggacagc tcccatcgaa 480
ctctcnaggc tctctanctc agggatgaag gaggtnaagt gatcgatnct cacaagcgan 540
                                                                   561
agetetegen enatatetge g
<210> 814
<211> 307
<212> DNA
<213> Homo sapiens
<220>
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<222> (1)...(307)
<223> n=A, T, C or G
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tttttttttt tttttggngg agggaaantt ncagacatag ctttattgct gactcctgcc 180
cccttcanag ccctagtcac aggcnncagg gntgttttgt aanttaaant ttcnggaaaa 240
tnggngtntt tntgcatnca anagaagggn tgccaaangn ggggtattgc ttctgggtgg 300
nttaccc
<210> 815
<211> 784
<212> DNA
<213> Homo sapiens
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<221> misc_feature
<222> (1)...(784)
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gttggattaa gctgcttatg agctctttga cagtgttgat tttgatcagt ggtttaaaaa 180
ccagcttctt ccagaattac aagtcattca caataggtat aagccattgc gacgcagggt 240
gatttggctc atcggtcagt ggatttctgt gaaattcaag tctgacttaa gacccatgct 300
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ttggaaacca tgttcacact actttttcag ttactgcagc aagttacaga atgtgacaca 480
aagatgcatg ttttgcatgt cctttcttgt gtgatcgaaa gagtcaacat gcagatacga 540
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aaac
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<211> 813
<212> DNA
<213> Homo sapiens
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<221> misc feature
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<223> n=A,T,C or G
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ageagetget geeagageee tettgtaget tetttatttt etgtttettt eeagetttee 180
taccetecta tecceeettg tgtttgggee acaattttga aataattttt attataggta 240
tgtgctgcca aagccagatt tttataaggt aaaataaatt aagaatttaa acagtaaaag 300
ccagtgtctc aaaatgtcag cattaaaatg tgaaggggac agcagggtgt gaaccggaaa 360
gcccttaagg tcaatgccag tgtccagacg agcagtgtag aaaagctccc tgtgtggttt 480
gtcgtgaggt ctgcttgtat ctcttcactg gcgttagttt cattagctct ttattctcct 540
tacgttcgag tgaatctgcc aagaacactg gtggatagta ttatcctaac acttttggtt 600
tgggggcggg gagggggcag ggaatagtga gctggcttta ccaccttcag gatctcgaat 660
tqqqcqcttq aacctaaqaa agattqtgga cttatcaaaa gtcaccgctc aqtqttcqtc 720
aagcatgtat ttatgtgacn atcatactag ggaggggatg gttgggaatt cttccatgtg 780
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                                                                 813
<210> 817
<211> 229
<212> DNA
<213> Homo sapiens
<220>
<221> misc feature
<222> (1)...(229)
<223> n=A,T,C or G
<400> 817
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acanacacat ttttttttcc aggtaaaagc tgtttttagt ttgtagtaca aatgtgactg 180
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<210> 818
<211> 781
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<212> DNA
<213> Homo sapiens
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tecteacate aegteetgee ceaggteact geataaataa gtgetttgga aagtatteat 180
ctagaaagta acataaatac tgtacataga aaagggttgc cgccccttag ccttcgcact 240
gececagaga getetecaca tattgeacac ggeetececa gecetgtggg gtecaggeet 300
ggctgtgtct ttggtagaag cttcagggac agttcctggg cagcccccac atctncaccc 360
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tggtggcctt ctaccangga tgctttcaca aggatgagac agaatcccaa tggtatgccc 480
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gnaanteete aaaneeetta eeaceaettg atgacenane atttttangg eetggettga 720
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а
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<211> 199
<212> DNA
<213> Homo sapiens
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<210> 820
<211> 211
<212> DNA
<213> Homo sapiens
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<221> misc_feature
<222> (1)...(211)
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<211> 264

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211
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<211> 952
<212> DNA
<213> Homo sapiens
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<221> misc feature
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<211> 587
<212> DNA
<213> Homo sapiens
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<223> n=A,T,C or G
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<210> 823
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<212> DNA
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<211> 520
<212> DNA
<213> Homo sapiens
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<223> n=A,T,C or G
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<211> 2064
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<213> Homo sapiens

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<213> Homo sapiens

<400> 827

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Arg Leu Met Asn Arg Asp Glu Asn Gly Gly Gly Ala Gly Gly Ser Gly

Ser His Gly Thr Leu Gly Leu Pro Ser Gly Gly Lys Cys Leu Leu Leu

Asp Cys Arg Pro Phe Leu Ala His Ser Ala Gly Tyr Ile Leu Gly Ser

Val Asn Val Arg Cys Asn Thr Ile Val Arg Arg Arg Ala Lys Gly Ser 70

Val Ser Leu Glu Gln Ile Leu Pro Ala Glu Glu Val Arg Ala Arg

Leu Arg Ser Gly Leu Tyr Ser Ala Val Ile Val Tyr Asp Glu Arg Ser 105

Pro Arg Ala Glu Ser Leu Arg Glu Asp Ser Thr Val Ser Leu Val Val 120

Gln Ala Leu Arg Arg Asn Ala Glu Arg Thr Asp Ile Cys Leu Leu Lys 135

Gly Gly Tyr Glu Arg Phe Ser Ser Glu Tyr Pro Glu Phe Cys Ser Lys 150 155

Thr Lys Ala Leu Ala Ala Ile Pro Pro Pro Val Pro Pro Ser Ala Thr 165

Glu Pro Leu Asp Leu Gly Cys Ser Ser Cys Gly Thr Pro Leu His Asp 180 185 190

Gln Gly	Gly Pro	Val Glu	Ile	Leu 200	Pro	Phe	Leu	Tyr	Leu 205	Gly	Ser	Ala
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Leu Leu 225	Asn Val	Ser Ser 230	_	Cys	Pro	Asn	His 235	Phe	Glu	Gly	His	Tyr 240
Gln Tyr	Lys Cys	Ile Pro 245	Val	Glu	Asp	Asn 250	His	Lys	Ala	Asp	Ile 255	Ser
Ser Trp	Phe Met 260	Glu Ala	Ile	Glu	Tyr 265	Ile	Asp	Ala	Val	Lys 270	Asp	Cys
Arg Gly	Arg Val 275	Leu Val	His	Cys 280	Gln	Ala	Gly	Ile	Ser 285	Arg	Ser	Ala
Thr Ile 290	-	Ala Tyr	Leu 295	Met	Met	Lys	Lys	Arg 300	Val	Arg	Leu	Glu
Glu Ala 305	Phe Glu	Phe Val	_	Gln	Arg	Arg	Ser 315	Ile	Ile	Ser	Pro	Asn 320
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Thr Ser	Cys Ala 340	Ala Glu	Ala	Ala	Ser 345	Pro	Ser	Gly	Pro	Leu 350	Arg	Glu
Arg Gly	Lys Thr 355	Pro Ala	Thr	Pro 360	Thr	Ser	Gln	Phe	Val 365	Phe	Ser	Phe
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